



RECEIVED

OCT 28 2002

TECH CENTER 1600/2900

FIG. 1-1

10	20	30	40	50	60	70
* *	* *	* *	* *	* *	* *	* *
GACGGATCGG	GAGATCTCCC	GATCCCCTAT	GGTCGACTCT	CAGTACAATC	TGCTCTGATG	CCGCATAGTT
80	90	100	110	120	130	140
* *	* *	* *	* *	* *	* *	* *
AAGCCAGTAT	CTGCTCCCTG	CTTGTGTGTT	GGAGGTCGCT	GAGTAGTGCG	CGAGCAAAAT	TTAAGCTACA
150	160	170	180	190	200	210
* *	* *	* *	* *	* *	* *	* *
ACAAGGCAAG	GCTTGACCGA	CAATTGCATG	AAGAATCTGC	TTAGGGTTAG	GCGTTTTGCG	CTGCTTCGCG
220	230	240	250	260	270	280
* *	* *	* *	* *	* *	* *	* *
ATGTACGGGC	CAGATATACG	CGTTGACATT	GATTATTGAC	TAGTTATTAA	TAGTAATCAA	TTACGGGGTC
290	300	310	320	330	340	350
* *	* *	* *	* *	* *	* *	* *
ATTAGTTCAT	AGCCCATATA	TGGAGTTCCG	CGTTACATAA	CTTACGGTAA	ATGGCCCCGC	TGGCTGACCG
360	370	380	390	400	410	420
* *	* *	* *	* *	* *	* *	* *
CCCAACGACC	CCCGCCATT	GACGTCAATA	ATGACGTATG	TTCCCATAGT	AACGCCAATA	GGGACTTTCC
430	440	450	460	470	480	490
* *	* *	* *	* *	* *	* *	* *
ATTGACGTCA	ATGGGTGGAC	TATTACGGT	AAACTGCCCA	CTTGGCAGTA	CATCAAGTGT	ATCATATGCC
500	510	520	530	540	550	560
* *	* *	* *	* *	* *	* *	* *
AAGTACGCCC	CCTATTGACG	TCAATGACGG	TAAATGGCCC	GCCTGGCATT	ATGCCAGTA	CATGACCTTA
570	580	590	600	610	620	630
* *	* *	* *	* *	* *	* *	* *
TGGGACTTTC	CTACTTGGCA	GTACATCTAC	GTATTAGTCA	TCGCTATTAC	CATGGTGATG	CGGTTTTGGC
640	650	660	670	680	690	700
* *	* *	* *	* *	* *	* *	* *
AGTACATCAA	TGGGCGTGGA	TAGCGGTTTG	ACTCACGGGG	ATTTCOAAGT	CTCCACCCCA	TTGACGTCAA
710	720	730	740	750	760	770
* *	* *	* *	* *	* *	* *	* *
TGGGAGTTTG	TTTTGGCACC	AAAATCAACG	GGACTTTCCA	AAATGTCGTA	ACAACTCCGC	CCCATTGACG
780	790	800	810	820	830	840
* *	* *	* *	* *	* *	* *	* *
CAAATGGGCG	GTAGGCGTGT	ACGGTGGGAG	GTCTATATAA	GCAGAGCTCT	CTGGCTAACT	AGAGAACCCA
850	860	870	880	890	900	910
* *	* *	* *	* *	* *	* *	* *
CTGCTTAACT	GGCTTATCGA	AATTAATACG	ACTCACTATA	GGGAGACCCA	AGCTTCGCAG	AATTCCTGCG
920	930	940	950	960	970	980
* *	* *	* *	* *	* *	* *	* *
GCTGCTACAG	TGTGTCCAGC	GTCCTGCCTG	GCTGTGCTGA	GCGCTGGAAC	AGTGGCGCAT	CATTCAAGTG
990	1000	1010	1020	1030	1040	1050
* *	* *	* *	* *	* *	* *	* *
CACAGTTACC	CATCCTGAGT	CTGGCACCTT	AACTGGCACA	ATTGCCAAAG	TCACAGGTGA	GCTCAGATGC
1060	1070	1080	1090	1100	1110	1120
* *	* *	* *	* *	* *	* *	* *
ATACCAGGAC	ATTGTATGAC	GTTCCCTGCT	CACATGCCTG	CTTTCTTCCT	ATAATACAGA	TGCTCAACTA
1130	1140	1150	1160	1170	1180	1190
* *	* *	* *	* *	* *	* *	* *
ACTGCTCATG	TCCTTATATC	ACAGAGGGAA	ATTGGAGCTA	TCTGAGGAAC	TGCCAGAAG	GGAAGGGCAG

FIG. 1-2

1200	1210	1220	1230	1240	1250	1260
* *	* *	* *	* *	* *	* *	* *
AGGGGTCTTG	CTCTCCTTGT	CTGAGCCATA	ACTCTTCTTT	CTACCTTCCA	GTGAACACCT	TCCCACCCCA
1270	1280	1290	1300	1310	1320	1330
* *	* *	* *	* *	* *	* *	* *
GGTCCACCTG	CTACCGCCGC	CGTCGGAGGA	GCTGGCCCTG	AATGAGCTCT	TGTCCCTGAC	ATGCCTGGTG
1340	1350	1360	1370	1380	1390	1400
* *	* *	* *	* *	* *	* *	* *
CGAGCTTTCA	ACCCTAAAGA	AGTGCTGGTG	CGATGGCTGC	ATGGAAATGA	GGAGCTGTCC	CCAGAAAGCT
1410	1420	1430	1440	1450	1460	1470
* *	* *	* *	* *	* *	* *	* *
ACCTAGTGTT	TGAGCCCCTA	AAGGAGCCAG	GCGAGGGAGC	CACCACCTAC	CTGGTGACAA	GCGTGTTCGC
1480	1490	1500	1510	1520	1530	1540
* *	* *	* *	* *	* *	* *	* *
TGTATCAGCT	GAAAGCTTGA	TATCGAATTC	CGGAGGCGGA	ACCGGCAGTG	CAGCCCGAAG	CCCCGCAGTC
1550	1560	1570	1580	1590		
* *	* *	* *	* *	* *		
CCCGAGCACG	CGTGGCC	ATG	CGT	CCC	CTG	CGC
		Met	Arg	Pro	Leu	Arg
				Pro	Arg	Ala
					Ala	Ala
					Leu	Leu
					Ala	Leu
					Leu	Leu
1600	1610	1620	1630	1640	1650	
* *	* *	* *	* *	* *	* *	
GCC	TCG	CTC	CTG	GCC	GCG	CCC
Ala	Ser	Leu	Leu	Ala	Ala	Pro
1660	1670	1680	1690	1700	1710	
* *	* *	* *	* *	* *	* *	
GTG	GAC	GCG	GCC	CGC	GCG	CTG
Val	Asp	Ala	Ala	Arg	Ala	Leu
1720	1730	1740	1750	1760	1770	
* *	* *	* *	* *	* *	* *	
TGC	CCC	CCG	CTG	CCA	CAC	AGC
Cys	Pro	Pro	Leu	Pro	His	Ser
1780	1790	1800	1810	1820		
* *	* *	* *	* *	* *		
CTC	AAC	CTC	GCC	TAT	GTG	GGC
Leu	Asn	Leu	Ala	Tyr	Val	Gly
1830	1840	1850	1860	1870	1880	
* *	* *	* *	* *	* *	* *	
CAC	TGG	CTG	CTG	GAG	CTT	GTC
His	Trp	Leu	Leu	Glu	Leu	Val
1890	1900	1910	1920	1930	1940	
* *	* *	* *	* *	* *	* *	
AAC	TTC	ACC	CAC	CTG	GAC	GGG
Asn	Phe	Thr	His	Leu	Asp	Gly
1950	1960	1970	1980	1990		
* *	* *	* *	* *	* *		
GGG	TTT	GAG	CTG	ATG	GGC	AGC
Gly	Phe	Glu	Leu	Met	Gly	Ser

FIG. 1-3

2000	2010	2020	2030	2040	2050		
* CAG GTG TTT GAG TGG AAG GAC TTG GTC TCC AGC CTG GCC AGG AGA TAC ATC GGT AGG	* Gln Val Phe Glu Trp Lys Asp Leu Val Ser Ser Leu Ala Arg Arg Tyr Ile Gly Arg	* 2060	* 2070	* 2080	* 2090	* 2100	* 2110
TAC GGA CTG GCG CAT GTT TCC AAG TGG AAC TTC GAG ACG TGG AAT GAG CCA GAC CAC	Tyr Gly Leu Ala His Val Ser Lys Trp Asn Phe Glu Thr Trp Asn Glu Pro Asp His	2120	2130	2140	2150	2160	
CAC GAC TTT GAC AAC GTC TCC ATG ACC ATG CAA GGC TTC CTG AAC TAC TAC GAT GCC	His Asp Phe Asp Asn Val Ser Met Thr Met Gln Gly Phe Leu Asn Tyr Tyr Asp Ala	2170	2180	2190	2200	2210	2220
TGC TCG GAG GGT CTG CGC GCC GCC AGC CCC GCC CTG CGG CTG GGA GGC CCC GGC GAC	Cys Ser Glu Gly Leu Arg Ala Ala Ser Pro Ala Leu Arg Leu Gly Gly Pro Gly Asp	2230	2240	2250	2260	2270	2280
TCC TTC CAC ACC CCA CCG CGA TCC CCG CTG AGC TGG GGC CTC CTG CGC CAC TGC CAC	Ser Phe His Thr Pro Pro Arg Ser Pro Leu Ser Trp Gly Leu Leu Arg His Cys His	2290	2300	2310	2320	2330	2340
GAC GGT ACC AAC TTC TTC ACT GGG GAG GCG GGC GTG CGG CTG GAC TAC ATC TCC CTC	Asp Gly Thr Asn Phe Phe Thr Gly Glu Ala Gly Val Arg Leu Asp Tyr Ile Ser Leu	2350	2360	2370	2380	2390	
CAC AGG AAG GGT GCG CGC AGC TCC ATC TCC ATC CTG GAG CAG GAG AAG GTC GTC GCG	His Arg Lys Gly Ala Arg Ser Ser Ile Ser Ile Leu Glu Gln Glu Lys Val Val Ala	2400	2410	2420	2430	2440	2450
CAG CAG ATC CGG CAG CTC TTC CCC AAG TTC GCG GAC ACC CCC ATT TAC AAC GAC GAG	Gln Gln Ile Arg Gln Leu Phe Pro Lys Phe Ala Asp Thr Pro Ile Tyr Asn Asp Glu	2460	2470	2480	2490	2500	2510
GCG GAC CCG CTG GTG GGC TGG TCC CTG CCA CAG CCG TGG AGG GCG GAC GTG ACC TAC	Ala Asp Pro Leu Val Gly Trp Ser Leu Pro Gln Pro Trp Arg Ala Asp Val Thr Tyr	2520	2530	2540	2550	2560	
GCG GCC ATG GTG GTG AAG GTC ATC GCG CAG CAT CAG AAC CTG CTA CTG GCC AAC ACC	Ala Ala Met Val Val Lys Val Ile Ala Gln His Gln Asn Leu Leu Leu Ala Asn Thr	2570	2580	2590	2600	2610	2620
ACC TCC GCC TTC CCC TAC GCG CTC CTG AGC AAC GAC AAT GCC TTC CTG AGC TAC CAC	Thr Ser Ala Phe Pro Tyr Ala Leu Leu Ser Asn Asp Asn Ala Phe Leu Ser Tyr His	2630	2640	2650	2660	2670	2680
CCG CAC CCC TTC GCG CAG CGC ACG CTC ACC GCG CGC TTC CAG GTC AAC AAC ACC CGC	Pro His Pro Phe Ala Gln Arg Thr Leu Thr Ala Arg Phe Gln Val Asn Asn Thr Arg						

	2690				2700				2710				2720				2730			
	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*			
	CCG	CCG	CAC	GTG	CAG	CTG	TTG	CGC	AAG	CCG	GTG	CTC	ACG	GCC	ATG	GGG	CTG	CTG	GCG	
	Pro	Pro	His	Val	Gln	Leu	Leu	Arg	Lys	Pro	Val	Leu	Thr	Ala	Met	Gly	Leu	Leu	Ala	
2740				2750			2760				2770				2780			2790		
	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	
	CTG	CTG	GAT	GAG	GAG	CAG	CTC	TGG	GCC	GAA	GTG	TCG	CAG	GCC	GGG	ACC	GTC	CTG	GAC	
	Leu	Leu	Asp	Glu	Glu	Gln	Leu	Trp	Ala	Glu	Val	Ser	Gln	Ala	Gly	Thr	Val	Leu	Asp	
	2800			2810			2820				2830				2840			2850		
	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	
	AGC	AAC	CAC	ACG	GTG	GGC	GTC	CTG	GCC	AGC	GCC	CAC	CGC	CCC	CAG	GGC	CCG	GCC	GAC	
	Ser	Asn	His	Thr	Val	Gly	Val	Leu	Ala	Ser	Ala	His	Arg	Pro	Gln	Gly	Pro	Ala	Asp	
	2860			2870			2880				2890				2900			2910		
	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	
	GCC	TGG	CGC	GCC	GCG	GTG	CTG	ATC	TAC	GCG	AGC	GAC	GAC	ACC	CGC	GCC	CAC	CCC	AAC	
	Ala	Trp	Arg	Ala	Ala	Val	Leu	Ile	Tyr	Ala	Ser	Asp	Asp	Thr	Arg	Ala	His	Pro	Asn	
	2920			2930			2940				2950				2960					
	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	
	CGC	AGC	GTC	GCG	GTG	ACC	CTG	CGG	CTG	CGC	GGG	GTG	CCC	CCC	GGC	CCG	GGC	CTG	GTC	
	Arg	Ser	Val	Ala	Val	Thr	Leu	Arg	Leu	Arg	Gly	Val	Pro	Pro	Gly	Pro	Gly	Leu	Val	
2970				2980			2990				3000				3010			3020		
	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	
	TAC	GTC	ACG	CGC	TAC	CTG	GAC	AAC	GGG	CTC	TGC	AGC	CCC	GAC	GGC	GAG	TGG	CGG	CGC	
	Tyr	Val	Thr	Arg	Tyr	Leu	Asp	Asn	Gly	Leu	Cys	Ser	Pro	Asp	Gly	Glu	Trp	Arg	Arg	
	3030			3040			3050				3060				3070			3080		
	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	
	CTG	GGC	CGG	CCC	GTC	TTC	CCC	ACG	GCA	GAG	CAG	TTC	CGG	CGC	ATG	CGC	GCG	GCT	GAG	
	Leu	Gly	Arg	Pro	Val	Phe	Pro	Thr	Ala	Glu	Gln	Phe	Arg	Arg	Met	Arg	Ala	Ala	Glu	
	3090			3100			3110				3120				3130					
	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	
	GAC	CCG	GTG	GCC	GCG	GCG	CCC	CGC	CCC	TTA	CCC	GCC	GGC	GGC	CGC	CTG	ACC	CTG	CGC	
	Asp	Pro	Val	Ala	Ala	Ala	Pro	Arg	Pro	Leu	Pro	Ala	Gly	Gly	Arg	Leu	Thr	Leu	Arg	
3140			3150				3160				3170				3180			3190		
	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	*	
	CCC	GCG	CTG	CGG	CTG	CCG	TCG	CTT	TTG	CTG	GTG	CAC	GTG	TGT	GCG	CGC	CCC	GAG	AAG	
	Pro	Ala	Leu	Arg	Leu	Pro	Ser	Leu	Leu	Leu	Val</									

FIG. 1-5

3370	3380	3390	3400	3410	3420
* * *	* * *	* * *	* * *	* * *	* * *
CTC TTT GTG TTC AGC CCA GAC ACA GGT GCT GTC TCT GGC TCC TAC CGA GTT CGA GCC					
Leu Phe Val Phe Ser Pro Asp Thr Gly Ala Val Ser Gly Ser Tyr Arg Val Arg Ala					
3430	3440	3450	3460	3470	3480
* * *	* * *	* * *	* * *	* * *	* * *
CTG GAC TAC TGG GCC CGA CCA GGC CCC TTC TCG GAC CCT GTG CCG TAC CTG GAG GTC					
Leu Asp Tyr Trp Ala Arg Pro Gly Pro Phe Ser Asp Pro Val Pro Tyr Leu Glu Val					
3490	3500	3510	3520	3530	3540
* * *	* * *	* * *	* * *	* * *	* * *
CCT GTG CCA AGA GGG CCC CCA TCC CCG GGC AAT CCA TGAG CCTGTGCTGA GCCCCAGTGG					
Pro Val Pro Arg Gly Pro Pro Ser Pro Gly Asn Pro					
3550	3560	3570	3580	3590	3600
* * *	* * *	* * *	* * *	* * *	* * *
GTTGCACCTC CACCGGCAGT CAGCGAGCTG GGGCTGCACT GTGCCCATGC TGCCCTCCCA TCACCCCTT					
3620	3630	3640	3650	3660	3670
* * *	* * *	* * *	* * *	* * *	* * *
TGCAATATAT TTTTATATTT TAAAAAAAAA AAAAAAAAAA AAAAAAAAAA AAAAAAAAAA AAAAAAAAAA					
3690	3700	3710	3720	3730	3740
* * *	* * *	* * *	* * *	* * *	* * *
AAAAAAAAAA AAAAAAAAAAG AATTCCTGCA GCCCAGGGGA TCCACTAGTT CTAGAGGGCC CGTTTAAACC					
3760	3770	3780	3790	3800	3810
* * *	* * *	* * *	* * *	* * *	* * *
CGCTGATCAG CCTCGACTGT GCCTTCTAGT TGCCAGCCAT CTGTTGTTTG CCCCTCCCC GTGCCTTCTT					
3830	3840	3850	3860	3870	3880
* * *	* * *	* * *	* * *	* * *	* * *
TGACCCTGGA AGGTGCCACT CCCACTGTCC TTTCTTAATA AAATGAGGAA ATTGCATCGC ATTGTCTGAG					
3900	3910	3920	3930	3940	3950
* * *	* * *	* * *	* * *	* * *	* * *
TAGGTGTCAT TCTATTCTGG GGGGTGGGGT GGGGCAGGAC AGCAAGGGGG AGGATTGGGA AGACAATAGC					
3970	3980	3990	4000	4010	4020
* * *	* * *	* * *	* * *	* * *	* * *
AGGCATGCTG GGGATGCGGT GGGCTCTATG GCTTCTGAGG CGGAAAGAAC CAGCTGGGGC TCGAGAGCTT					
4040	4050	4060	4070	4080	4090
* * *	* * *	* * *	* * *	* * *	* * *
GGCGTAATCA TGGTCATAGC TGTTCCTGT GTGAAATTGT TATCCGCTCA CAATTCCACA CAACATACGA					
4110	4120	4130	4140	4150	4160
* * *	* * *	* * *	* * *	* * *	* * *
GCCGGAAGCA TAAAGTGTA AGCCTGGGGT GCCTAATGAG TGAGCTAACT CACATTAATT GCGTTGCGCT					
4180	4190	4200	4210	4220	4230
* * *	* * *	* * *	* * *	* * *	* * *
CACTGCCCCG TTTCCAGTCG GGAAACCTGT CGTGCCAGCT GCATTAATGA ATCGGCCAAC GCGCGGGGAG					
4250	4260	4270	4280	4290	4300
* * *	* * *	* * *	* * *	* * *	* * *
AGGCGGTTTG CGTATTGGGC GCTCTTCCGC TTCCTCGCTC ACTGACTCGC TGCCTCGGT CGTTCGGCTG					
4320	4330	4340	4350	4360	4370
* * *	* * *	* * *	* * *	* * *	* * *
CGGCGAGCGG TATCAGCTCA CTCAAAGGCG GTAATACGGT TATCCACAGA ATCAGGGGAT AACGCAGGAA					

FIG. 1-6

4390	4400	4410	4420	4430	4440	4450
* *	* *	* *	* *	* *	* *	* *
AGAACATGTG	AGCAAAAGGC	CAGCAAAAGG	CCAGGAACCG	TAAAAAGGCC	GCGTTGCTGG	CGTTTTTCCA
4460	4470	4480	4490	4500	4510	4520
* *	* *	* *	* *	* *	* *	* *
TAGGCTCCGC	CCCCCTGACG	AGCATCACAA	AAATCGACGC	TCAAGTCAGA	GGTGGCGAAA	CCCGACAGGA
4530	4540	4550	4560	4570	4580	4590
* *	* *	* *	* *	* *	* *	* *
CTATAAAGAT	ACCAGGCGTT	TCCCCCTGGA	AGCTCCCTCG	TGCGCTCTCC	TGTTCCGACC	CTGCCGCTTA
4600	4610	4620	4630	4640	4650	4660
* *	* *	* *	* *	* *	* *	* *
CCGGATACCT	GTCCGCCTTT	CTCCCTTCGG	GAAGCGTGGC	GCTTTCTCAA	TGCTCACGCT	GTAGGTATCT
4670	4680	4690	4700	4710	4720	4730
* *	* *	* *	* *	* *	* *	* *
CAGTTCGGTG	TAGGTCGTTT	GCTCCAAGCT	GGGCTGTGTG	CACGAACCCC	CCGTTACGCC	CGACCGCTGC
4740	4750	4760	4770	4780	4790	4800
* *	* *	* *	* *	* *	* *	* *
GCCTTATCCG	GTAACATATG	TCTTGAGTCC	AACCCGGTAA	GACACGACTT	ATCGCCACTG	GCAGCAGCCA
4810	4820	4830	4840	4850	4860	4870
* *	* *	* *	* *	* *	* *	* *
CTGGTAACAG	GATTAGCAGA	GCGAGGTATG	TAGGCGGTGC	TACAGAGTTC	TTGAAGTGGT	GGCCTAACTA
4880	4890	4900	4910	4920	4930	4940
* *	* *	* *	* *	* *	* *	* *
CGGCTACACT	AGAAGGACAG	TATTTGGTAT	CTGCGCTCTG	CTGAAGCCAG	TTACCTTCGG	AAAAAGAGTT
4950	4960	4970	4980	4990	5000	5010
* *	* *	* *	* *	* *	* *	* *
GGTAGCTCTT	GATCCGGCAA	ACAAACCACC	GCTGGTAGCG	GTGGTTTTTT	TGTTTGCAAG	CAGCAGATTA
5020	5030	5040	5050	5060	5070	5080
* *	* *	* *	* *	* *	* *	* *
CGCGCAGAAA	AAAAGGATCT	CAAGAAGATC	CTTTGATCTT	TTCTACGGGG	TCTGACGCTC	AGTGAACGA
5090	5100	5110	5120	5130	5140	5150
* *	* *	* *	* *	* *	* *	* *
AAACTCACGT	TAAGGGATTT	TGGTCATGAG	ATTATCAAAA	AGGATCTTCA	CCTAGATCCT	TTTAAATTAA
5160	5170	5180	5190	5200	5210	5220
* *	* *	* *	* *	* *	* *	* *
AAATGAAGTT	TTAAATCAAT	CTAAAGTATA	TATGAGTAAA	CTTGGTCTGA	CAGTTACCAA	TGCTTAATCA
5230	5240	5250	5260	5270	5280	5290
* *	* *	* *	* *	* *	* *	* *
GTGAGGCACC	TATCTCAGCG	ATCTGTCTAT	TTCGTTTCATC	CATAGTTGCC	TGACTCCCCG	TCGTGTAGAT
5300	5310	5320	5330	5340	5350	5360
* *	* *	* *	* *	* *	* *	* *
AACTACGATA	CGGGAGGGCT	TACCATCTGG	CCCCAGTGCT	GCAATGATAC	CGCGAGACCC	ACGCTCACCG
5370	5380	5390	5400	5410	5420	5430
* *	* *	* *	* *	* *	* *	* *
GCTCCAGATT	TATCAGCAAT	AAACCAGCCA	GCCGGAAGGG	CCGAGCGCAG	AAGTGGTCCT	GCAACTTTAT
5440	5450	5460	5470	5480	5490	5500
* *	* *	* *	* *	* *	* *	* *
CCGCTCCAT	CCAGTCTATT	AATTGTTGCC	GGAAGCTAG	AGTAAGTAGT	TCGCCAGTTA	ATAGTTTGCG



FIG. 1-7

5510	5520	5530	5540	5550	5560	5570
* *	* *	* *	* *	* *	* *	* *
CAACGTTGTT	GCCATTGCTA	CAGGCATCGT	GGTGTACGCG	TCGTCGTTTG	GTATGGCTTC	ATTCAGCTCC
5580	5590	5600	5610	5620	5630	5640
* *	* *	* *	* *	* *	* *	* *
GGTTCCCAAC	GATCAAGGCG	AGTTACATGA	TCCCCCATGT	TGTGCAAAAA	AGCGGTTAGC	TCCTTCGGTC
5650	5660	5670	5680	5690	5700	5710
* *	* *	* *	* *	* *	* *	* *
CTCCGATCGT	TGTCAGAAGT	AAGTTGGCCG	CAGTGTTATC	ACTCATGGTT	ATGGCAGCAC	TGCATAATTC
5720	5730	5740	5750	5760	5770	5780
* *	* *	* *	* *	* *	* *	* *
TCTTACTGTC	ATGCCATCCG	TAAGATGCTT	TTCTGTGACT	GGTGAGTACT	CAACCAAGTC	ATTCTGAGAA
5790	5800	5810	5820	5830	5840	5850
* *	* *	* *	* *	* *	* *	* *
TAGTGTATGC	GGCGACCGAG	TTGCTCTTGC	CCGGCGTCAA	TACGGGATAA	TACCGCGCCA	CATAGCAGAA
5860	5870	5880	5890	5900	5910	5920
* *	* *	* *	* *	* *	* *	* *
CTTTAAAAGT	GCTCATCATT	GGAAAACGTT	CTTCGGGGCG	AAAACTCTCA	AGGATCTTAC	CGCTGTTGAG
5930	5940	5950	5960	5970	5980	5990
* *	* *	* *	* *	* *	* *	* *
ATCCAGTTTCG	ATGTAACCCA	CTCGTGCACC	CAACTGATCT	TCAGCATCTT	TTACTTTCAC	CAGCGTTTCT
6000	6010	6020	6030	6040	6050	6060
* *	* *	* *	* *	* *	* *	* *
GGGTGAGCAA	AAACAGGAAG	GCAAAATGCC	GCAAAAAAGG	GAATAAGGGC	GACACGGAAA	TGTTGAATAC
6070	6080	6090	6100	6110	6120	6130
* *	* *	* *	* *	* *	* *	* *
TCATACTCTT	CCTTTTTCAA	TATTATTGAA	GCATTTATCA	GGGTTATTGT	CTCATGAGCG	GATACATATT
6140	6150	6160	6170	6180	6190	6200
* *	* *	* *	* *	* *	* *	* *
TGAATGTATT	TAGAAAAATA	AACAAATAGG	GGTTCGCGC	ACATTTCCCC	GAAAAGTGCC	ACCTGACGTC

RECEIVED
OCT 23 2002
PATENT & TRADEMARK OFFICE

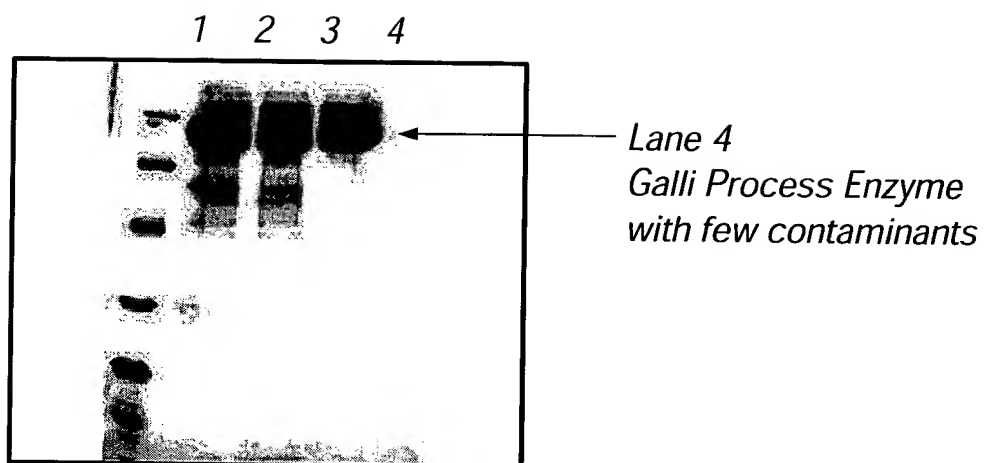
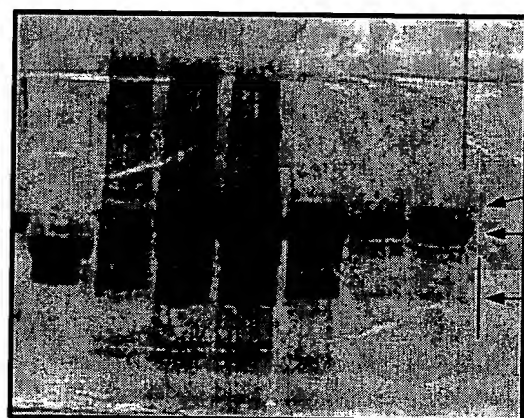


FIG. 2

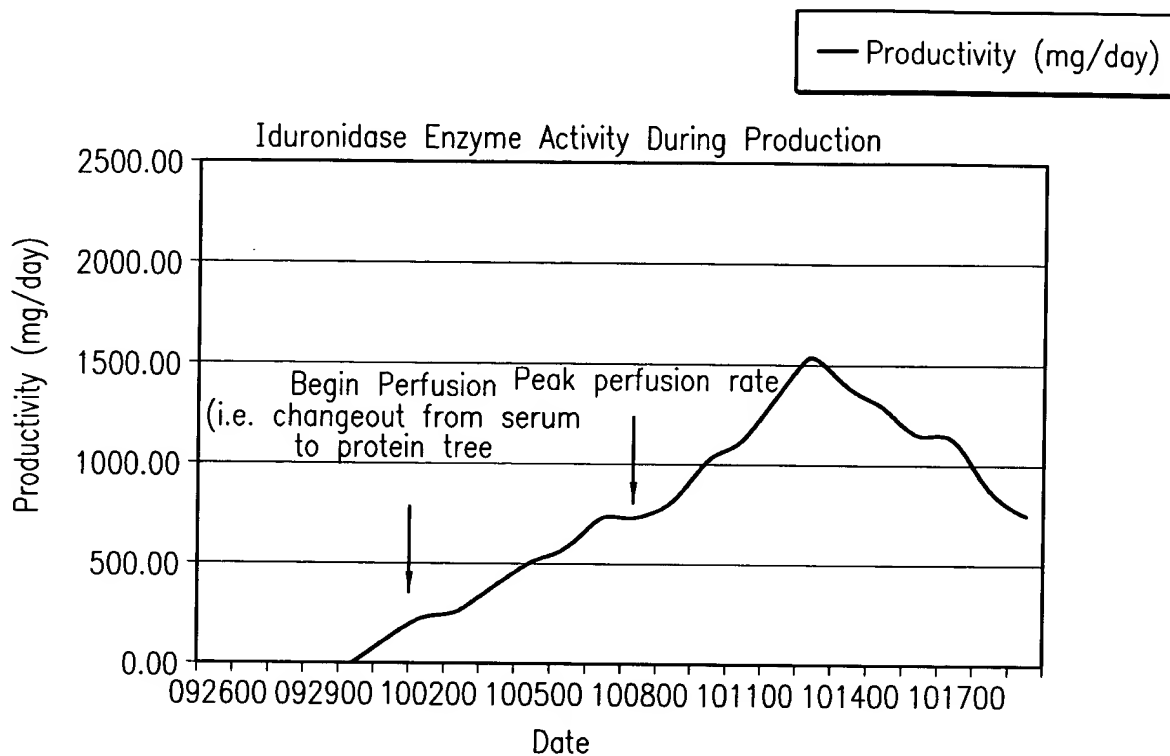


FIG. 3A

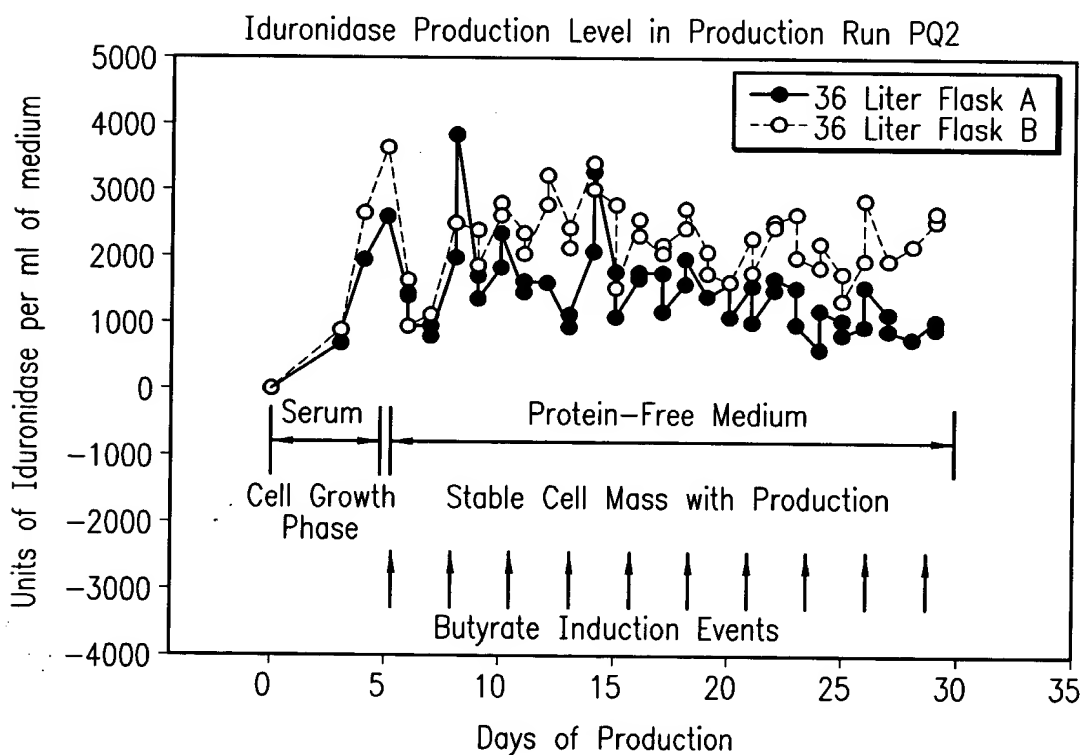
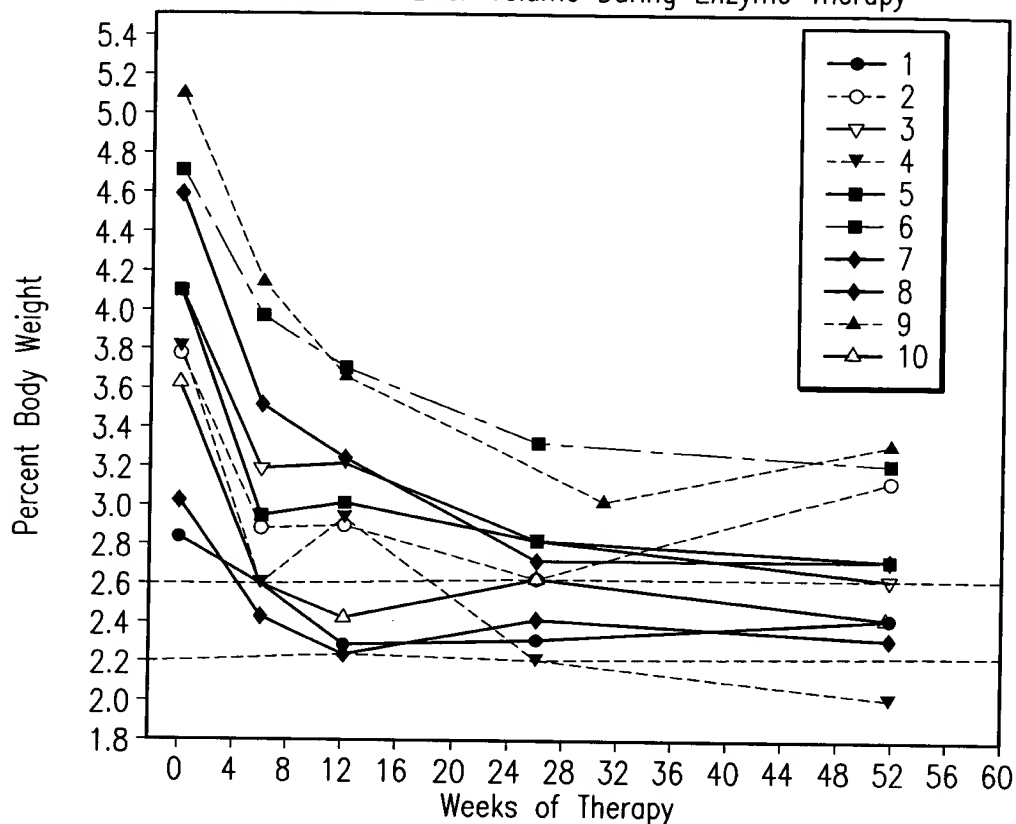


FIG. 3B

FIG. 4

Reduction in Liver Volume During Enzyme Therapy



Urinary GAG Excretion During Enzyme Therapy

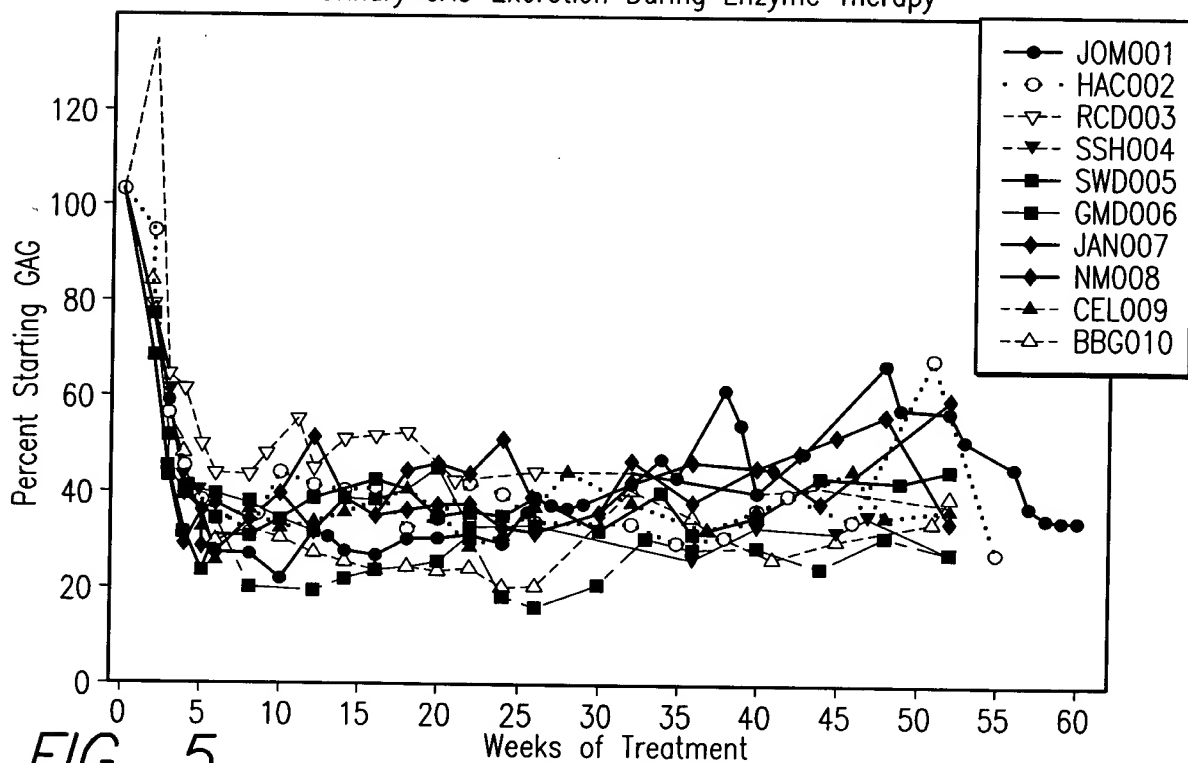
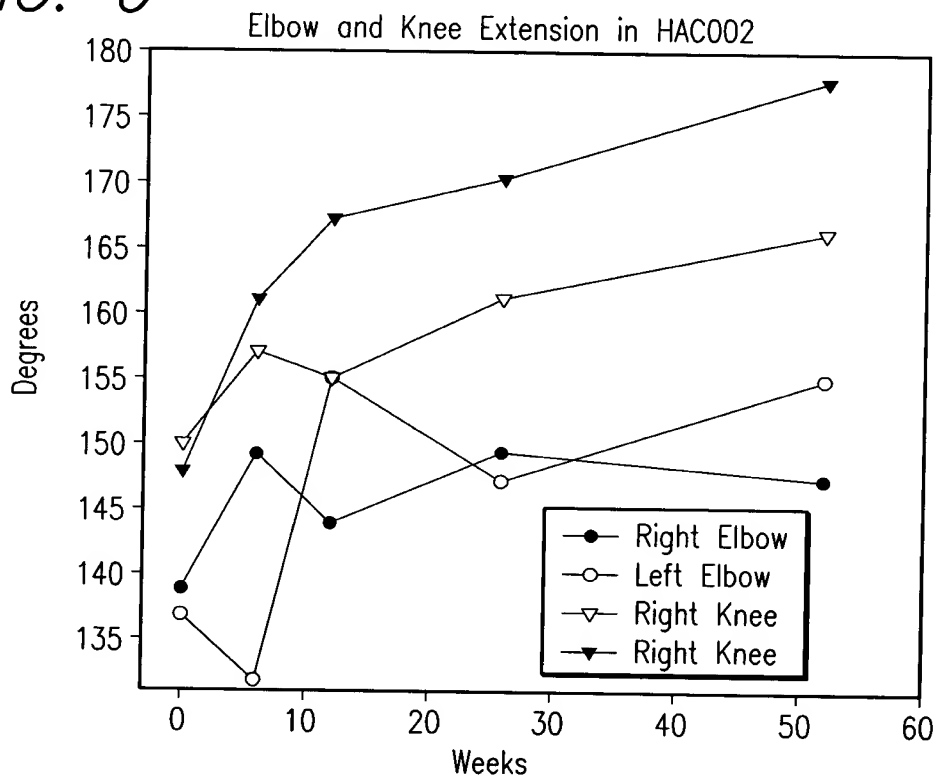


FIG. 5

FIG. 6



Shoulder flexion to 104 weeks in four patients with most restriction

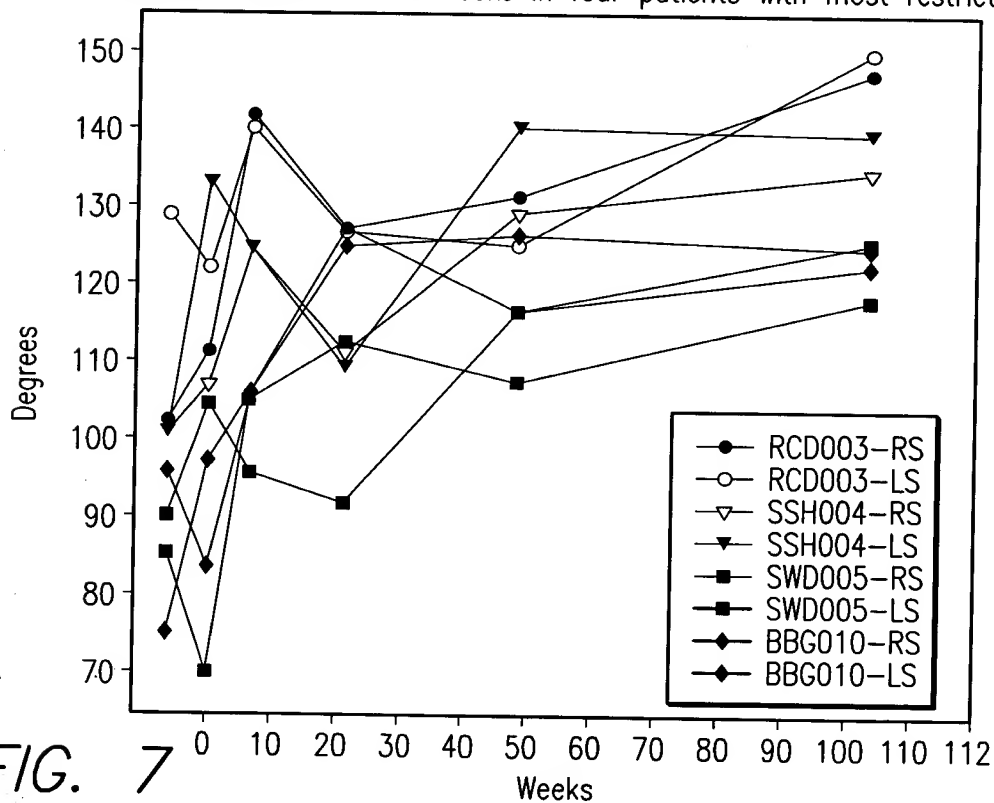


FIG. 7

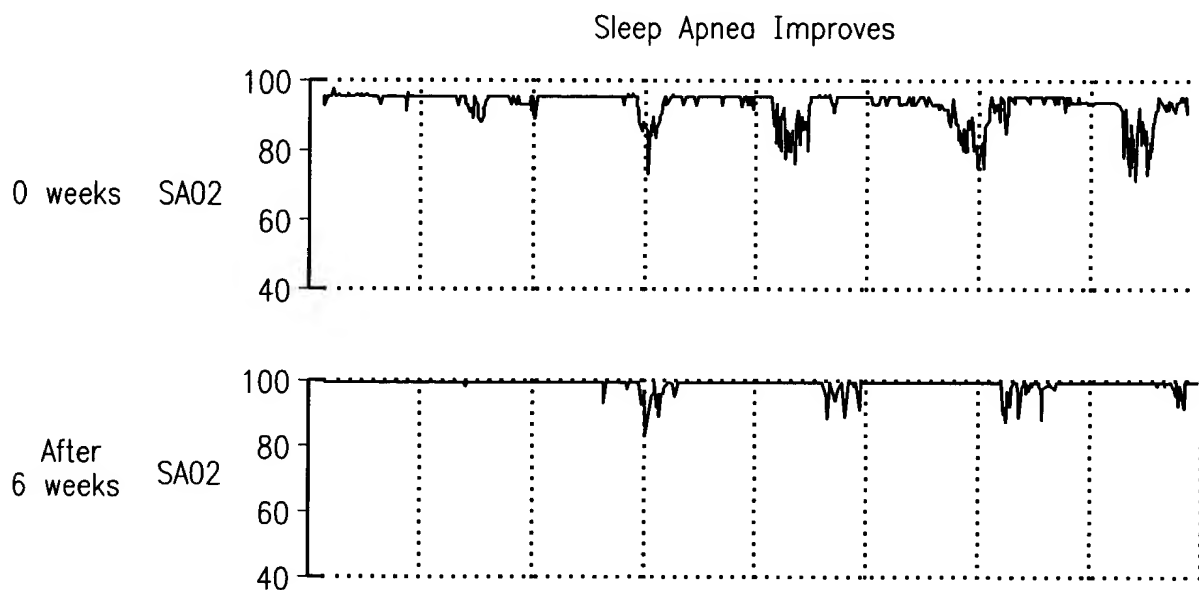


FIG. 8

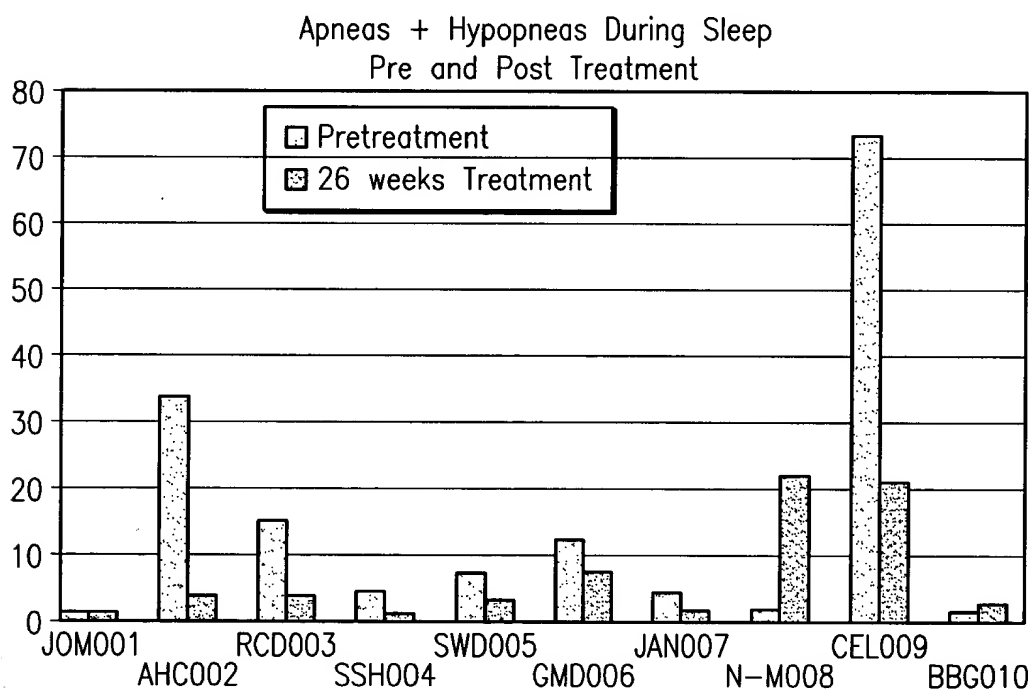


FIG. 9

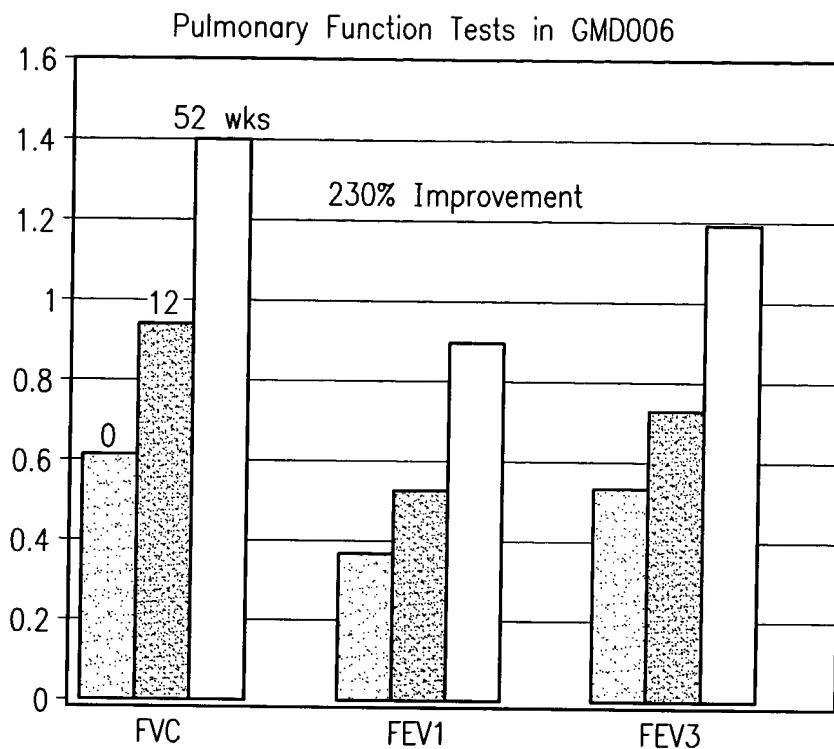


FIG. 10

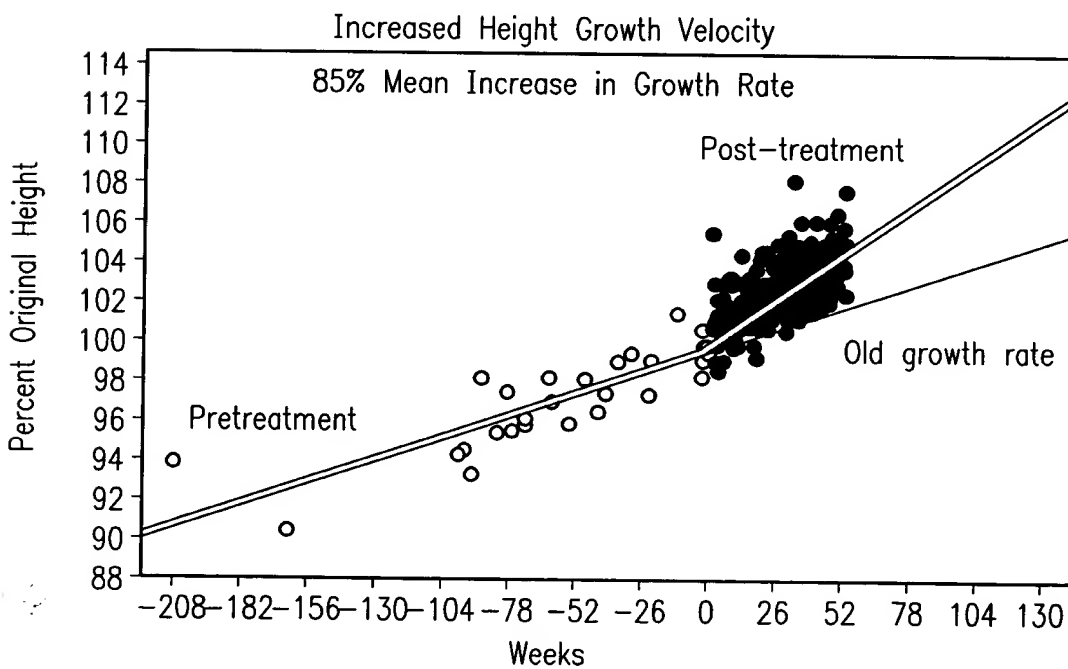


FIG. 11



FIG. 12

Chinese Hamster Ovary Host Protein Contamination by ELISA Assay

SOURCE AND BATCH NUMBER	CHOP PROTEIN CONTAMINATION (microgram per milligram)	PERCENT CHOP CONTAMINATION	PURITY OF THE ENZYME FROM CHOP
Prior Process (Carson/REI)			
C9002	14	1.4%	98.6%
C9003	24	2.4%	97.6%
C9004	16	1.6%	98.4%
New Process (Galli)			
P1003	<1.3	<0.13%	>99.9%
P1006	1.2	0.12%	99.9%
P1007	<0.6	<0.06%	>99.9%
P1008	<0.67	<0.067%	>99.9%

Comparison of Galli and Carson Material

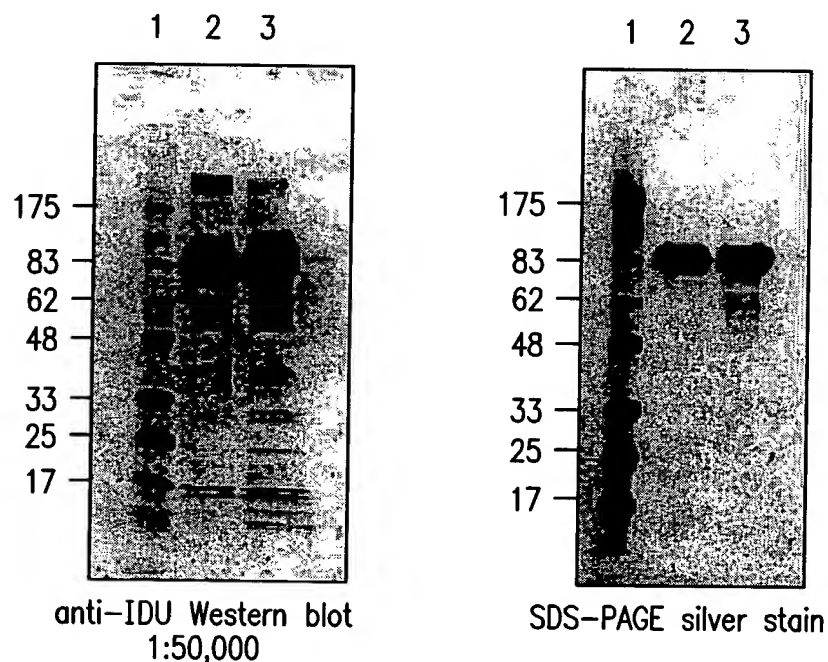


FIG. 13

1 Marker
2 Galli Referenced-0201
3 Carson C9002
5µg/lane